

EXPLOITING LIPID METABOLISM BY HSV-1: A CHALLENGE TO RETHINK NEW THERAPIES FOR ALZHEIMER'S DISEASE



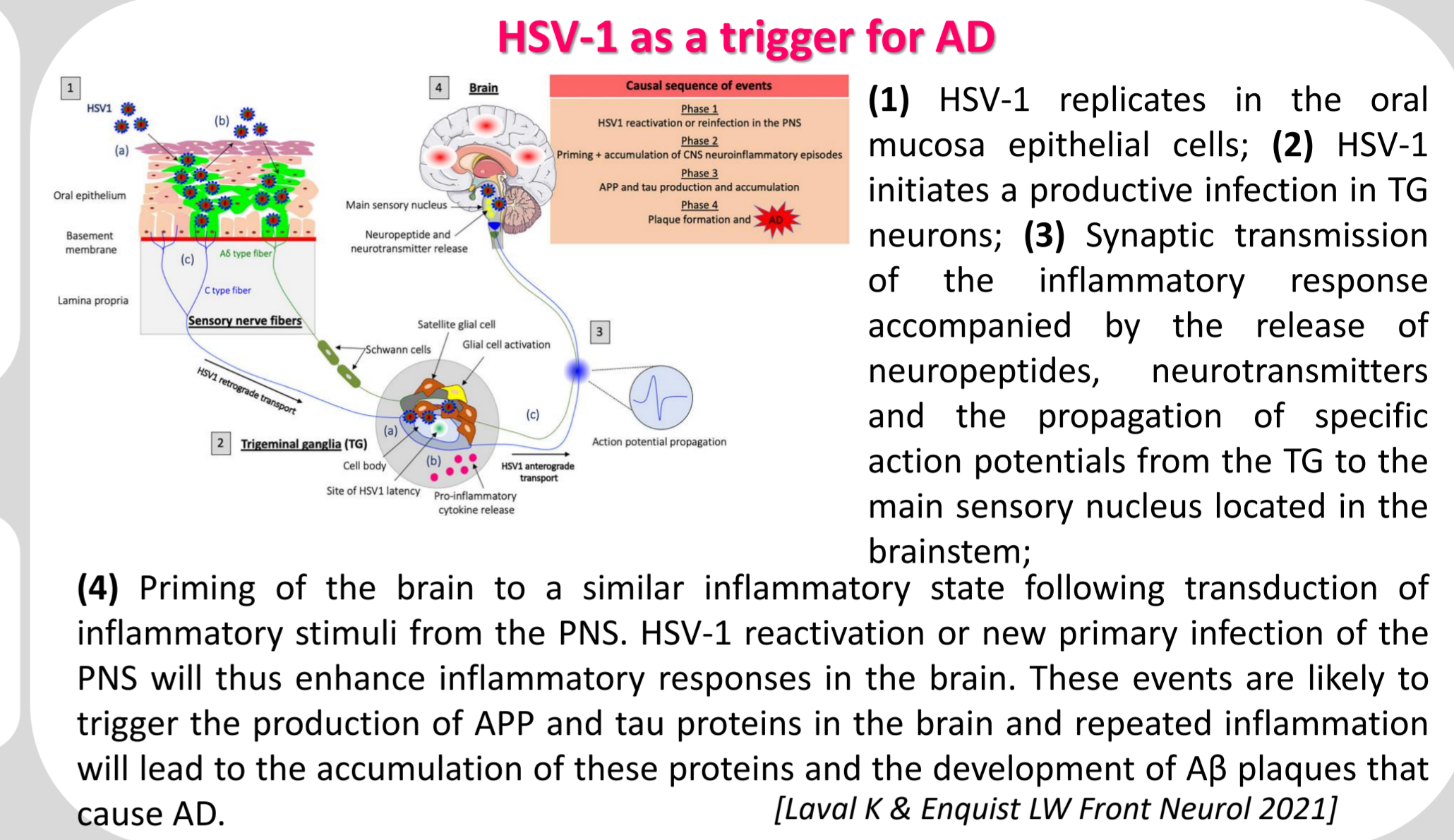
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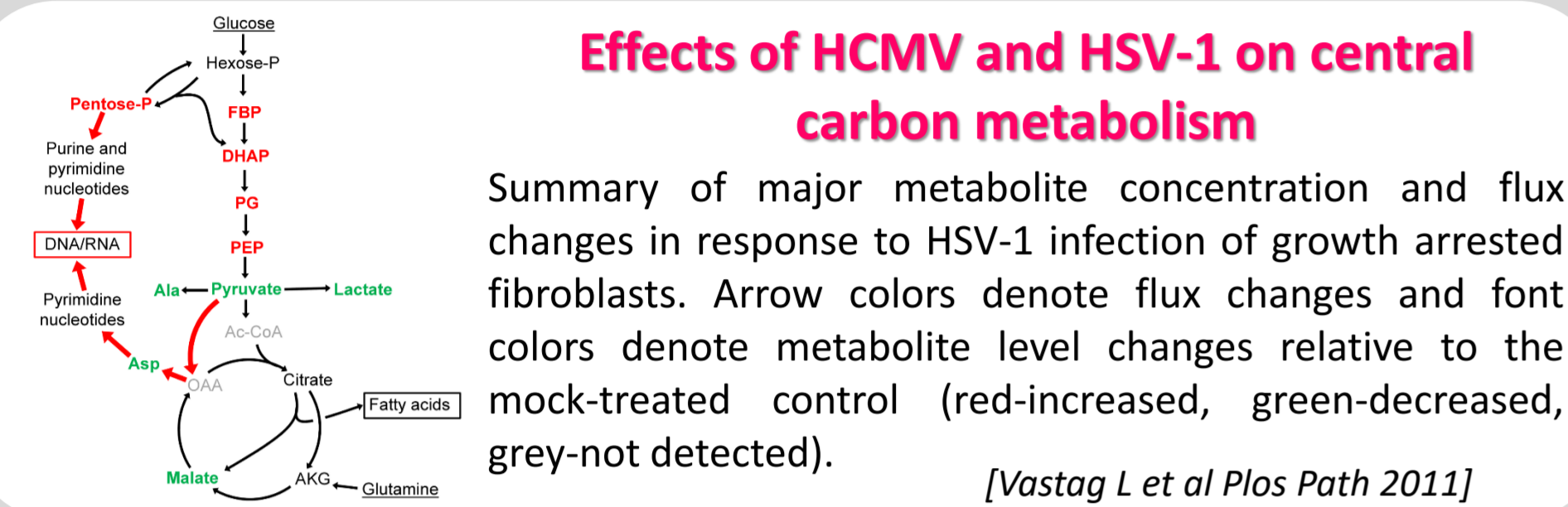
ABSTRACT

Herpes simplex virus-1 (HSV-1) establishes a life-long latent infection and can enter the brain via retrograde axonal transport. Recurrent reactivation of HSV-1 may lead to neurodegenerative disorders, including Alzheimer's disease (AD), although the underlying mechanisms have not been fully elucidated yet. Lipids constitute the bulk of the brain dry mass and alteration of lipid metabolism is a key component in AD. Considering that the mechanisms for remodeling of metabolism by HSV-1 are still poorly understood, we aim at dissecting the host metabolic pathways modulated by infection in a neuronal-like cell line to identify pathways that might be targeted to prevent AD. Specifically, we found an increase in both de novo synthesis and lipid storage following HSV-1 infection. In addition, anti-AD compounds targeting lipid metabolism (e.g. CMS121, C75) impaired HSV-1 infectivity. Overall, our data unveil new aspects of HSV-1-AD interplay and uncover new potential targets to rethink new possible therapies.

BACKGROUND



Effects of HCMV and HSV-1 on central carbon metabolism

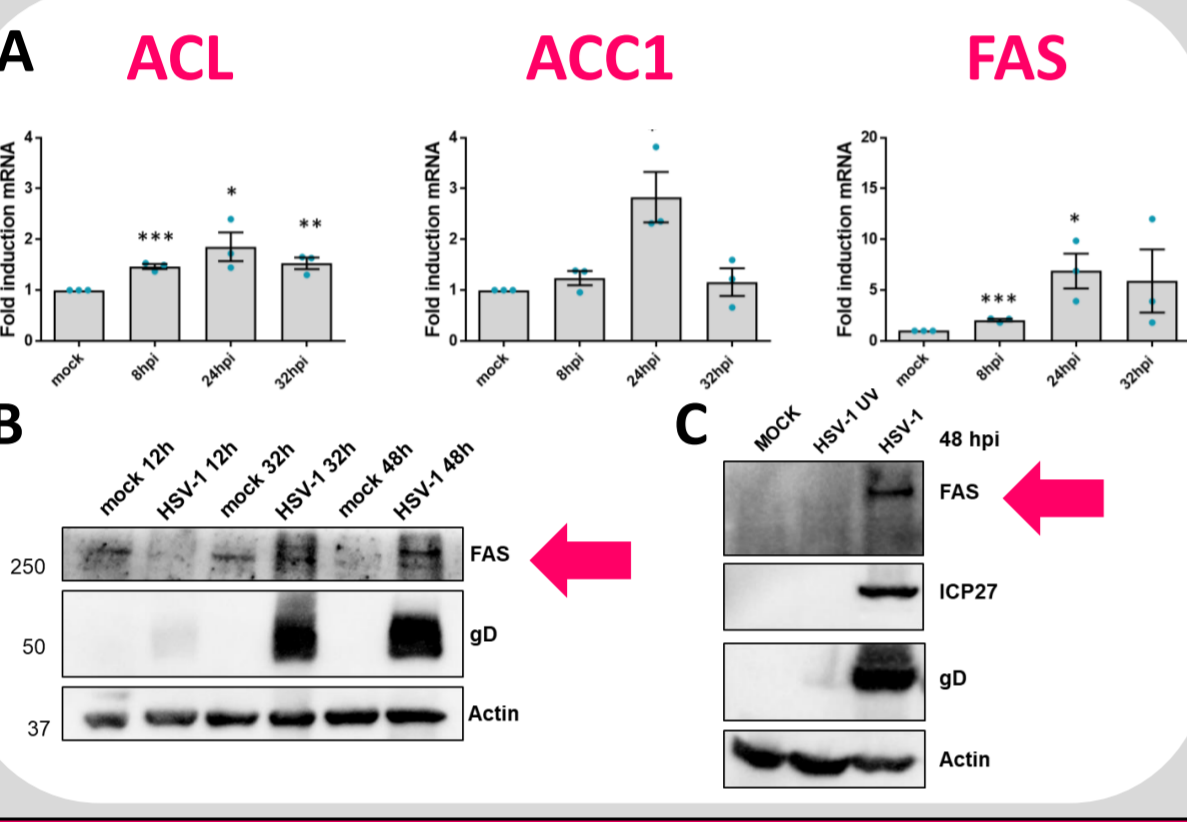


AIM OF THE PROJECT

Could anti-AD drugs modulate HSV-1 replication and lipid metabolism?

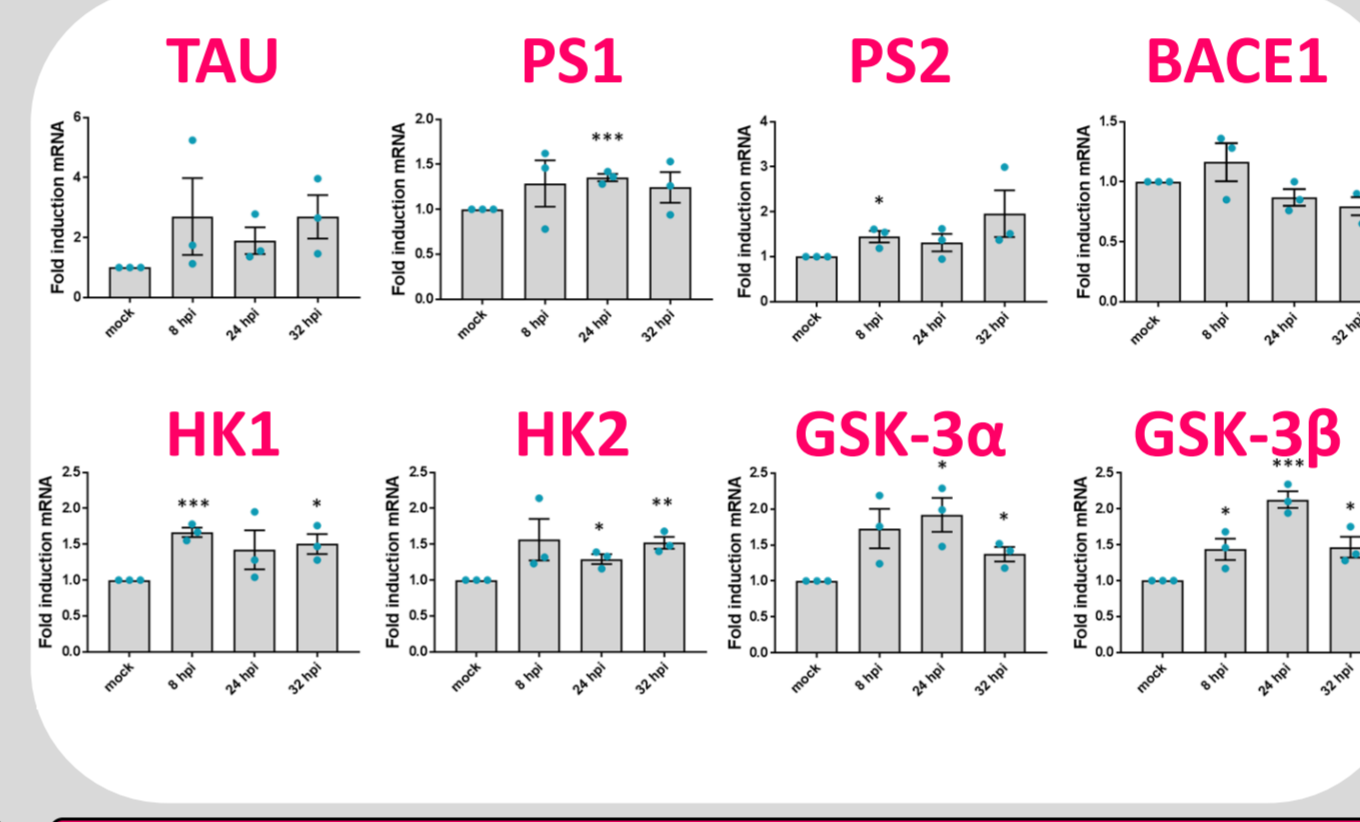
RESULTS

HSV-1 induces lipogenic enzymes



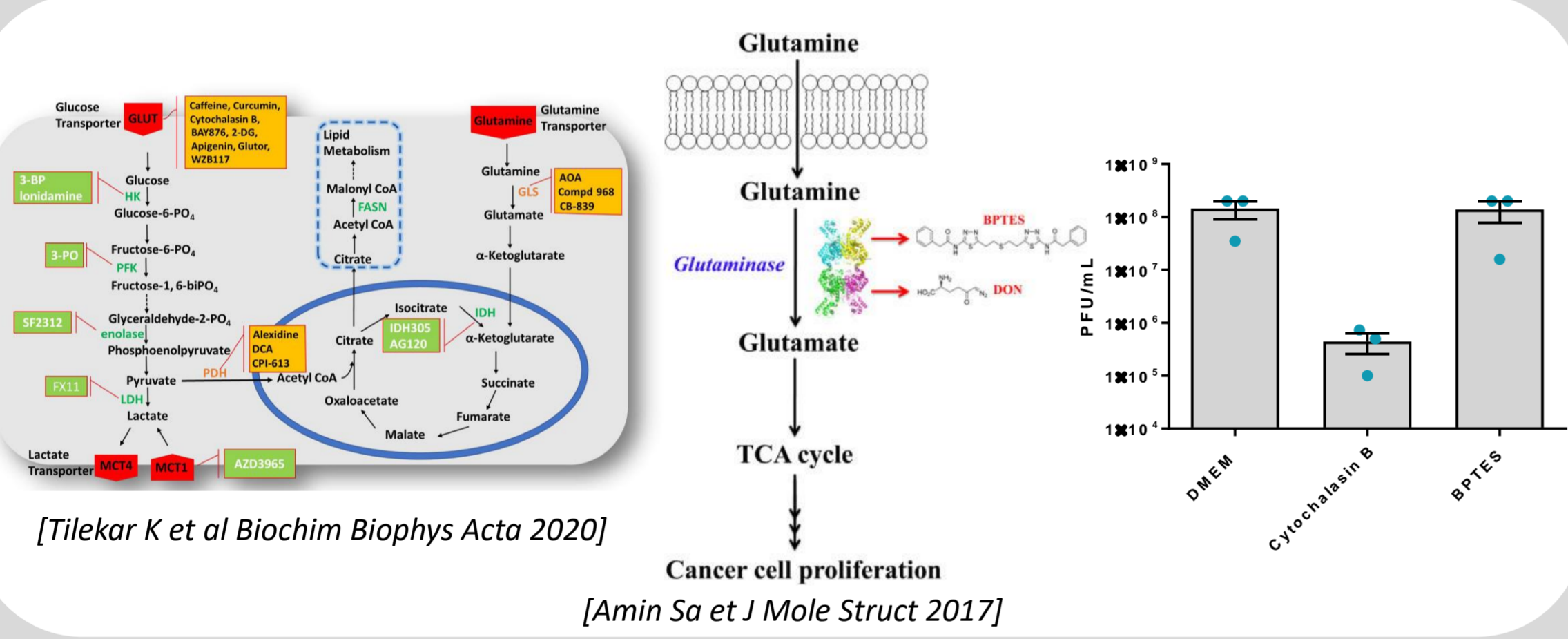
SH-SY5Y cells were infected with HSV-1 (MOI 1). (A) At 8, 24, and 32 hpi, total RNA was isolated and subjected to RT-qPCR to measure mRNA expression levels of the lipogenic enzymes ATP citrate lyase (ACL), Acetyl-CoA carboxylase (ACC1), and Fatty acid synthase (FAS). Values were normalized to the housekeeping glyceraldehyde-3-phosphate dehydrogenase (GAPDH) gene mRNA and plotted as fold induction relative to mock-infected cells (set at 1). The results were expressed as the mean ± standard error of the mean (SEM) of three independent experiments (***P* < 0.001; **P* < 0.01; **P* < 0.05; unpaired t test) for the comparison between mock- and infected cells. (B) Western blot analysis of protein lysates from mock- or infected cells using the indicated antibodies. A representative blot is shown. (C) SH-SY5Y cells were infected with HSV-1 or HSV-1 UV (MOI 1). Western blot analysis of protein lysates from mock- or infected cells using the indicated antibodies. A representative blot is shown.

Modulation of AD-related genes by HSV-1



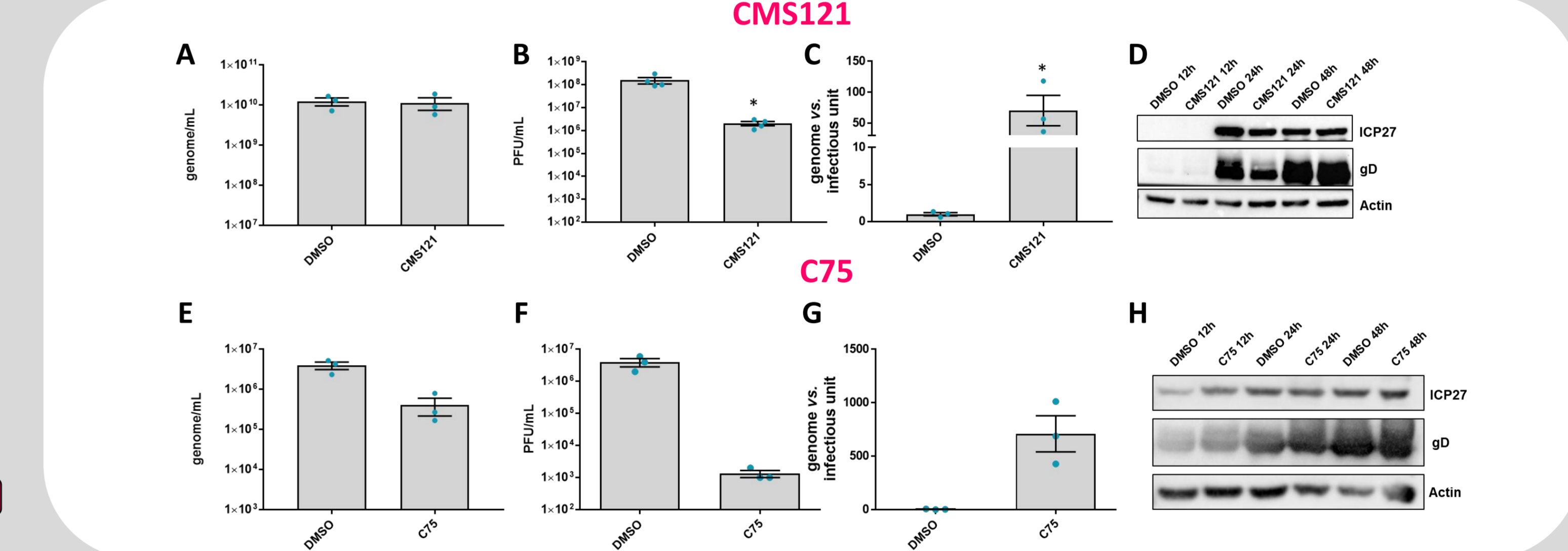
SH-SY5Y cells were infected with HSV-1 (MOI 1). At 8, 24, and 32 hpi, total RNA was isolated and subjected to RT-qPCR to measure mRNA expression levels of the AD-related genes: TAU, presenilin-1 (PS1), presenilin-2 (PS2), β-site APP cleaving enzyme 1 (BACE1), Hexokinase 1 (HK1), Hexokinase 2 (HK2), glycogen synthase kinase 3α (GSK-3α), glycogen synthase kinase 3β (GSK-3β). Values were normalized to the housekeeping glyceraldehyde-3-phosphate dehydrogenase (GAPDH) gene mRNA and plotted as fold induction relative to mock-infected cells (set at 1). The results were expressed as the mean ± standard error of the mean (SEM) of three independent experiments (***P* < 0.001; **P* < 0.01; **P* < 0.05; unpaired t test) for the comparison between mock- and infected cells.

Glucose is the main source of HSV-1 induced lipogenesis



SH-SY5Y cells were infected with HSV-1 (MOI 1) and treated with a non-toxic concentration of Cytochalasin B or BPTES (5 μM). The extent of HSV-1 replication was assessed by titrating the infectivity of supernatants and cell-associated viruses by standard plaque assay. The bars show means and SEMs from three independent experiments (unpaired t-test).

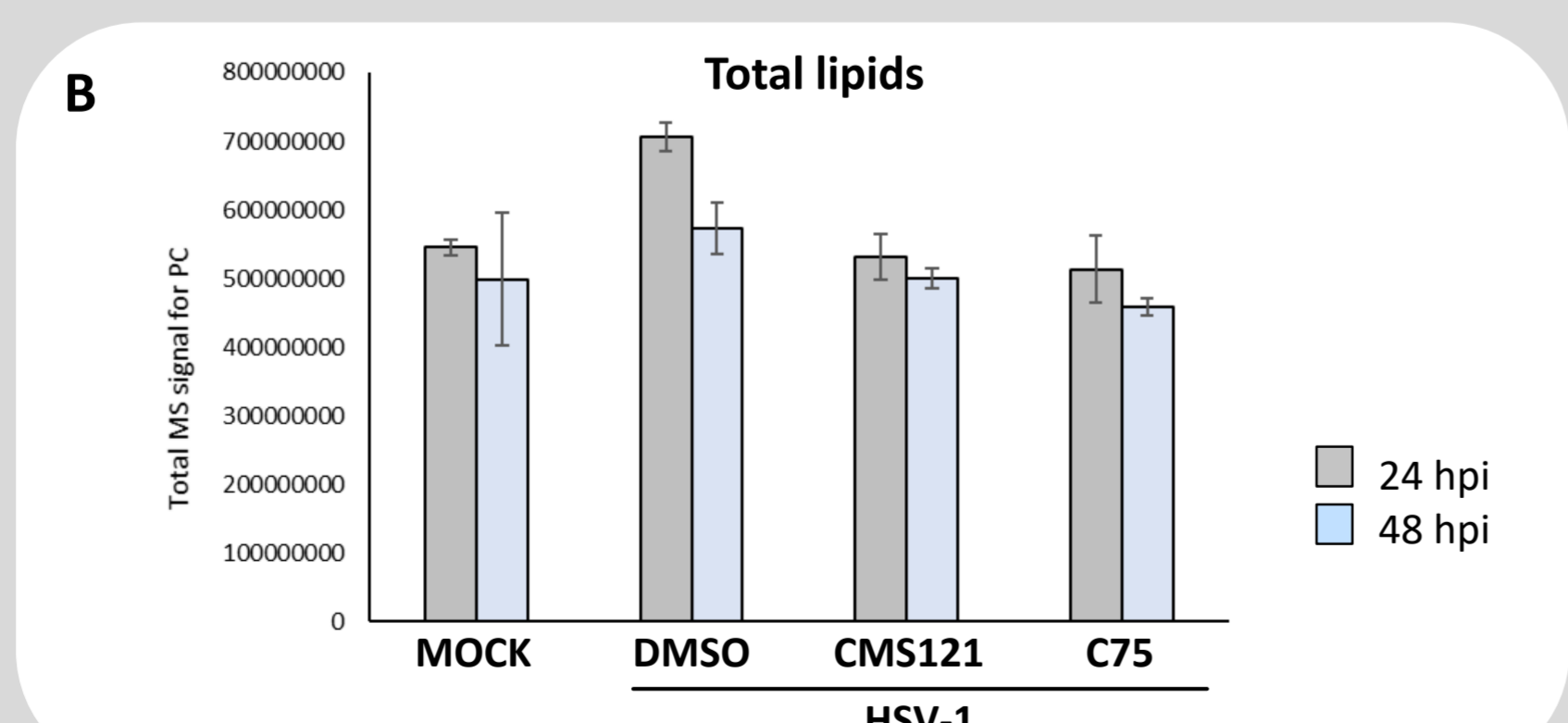
CMS121 and C75 reduce the infectivity of HSV-1 viral particles



HSV-1 induced lipogenesis

LIPID NAME	HSV-1/MOCK		CMS121/DMSO		C75/DMSO	
	24h	48h	24h	48h	24h	48h
EtherPC_300	1.08	1.38	1.37	1.35	1.27	1.08
EtherPC_301 iso 1	1.24	1.01	1.19	1.25	1.08	1.05
EtherPC_301 iso 2	1.04	1.00	1.01	1.15	0.81	0.77
EtherPC_320	1.12	1.41	0.99	0.99	1.02	1.05
EtherPC_340	1.23	1.52	1.18	1.19	0.91	1.21
EtherPC_341	1.06	1.05	1.12	1.23	1.07	0.95
EtherPC_342 iso 1	0.92	1.10	0.87	1.02	1.26	0.89
EtherPC_342 iso 2	1.18	1.09	0.90	1.03	0.64	0.99
EtherPC_361	1.28	1.12	1.61	1.65	0.85	0.77
EtherPC_362 iso 1	1.08	1.13	0.84	0.98	1.09	0.94
EtherPC_362 iso 2	1.12	1.17	1.17	1.10	0.96	0.98
EtherPC_363 iso 1	1.06	1.19	0.96	0.94	1.47	1.31
EtherPC_363 iso 2	1.13	1.08	0.85	0.92	0.69	0.98
EtherPC_382	1.12	1.32	1.12	1.11	0.97	0.93
EtherPC_383	1.02	1.23	0.95	0.91	1.49	1.42
EtherPC_384 iso 1	1.15	1.12	0.92	0.90	1.31	1.39
EtherPC_384 iso 2	1.02	1.14	0.98	0.97	1.05	0.97
EtherPC_385	1.06	1.20	0.97	0.95	1.38	1.24
EtherPC_403	1.20	1.56	1.12	1.11	1.21	1.27
EtherPC_404 iso 1	0.95	1.02	0.86	0.91	1.02	1.17
EtherPC_404 iso 2	1.03	1.19	1.03	1.02	1.22	1.17
EtherPC_404 iso 3	0.90	1.03	0.90	0.69	0.92	0.89
LPC_181	0.58	1.07	0.90	1.25	1.26	0.83
PC_280	0.91	0.71	1.03	1.05	1.17	1.32
PC_281 iso 1	0.84	0.71	0.70	0.87	1.12	1.47
PC_300_140_160	1.00	1.00	0.90	0.95	0.99	1.11
PC_301 iso 3	1.03	0.89	0.80	0.92	1.32	1.31
PC_310_150_160	1.07	1.07	0.80	0.96	0.98	1.08
PC_320_160_180	0.92	1.02	0.78	0.86	1.02	1.17
PC_321 iso 1	0.94	0.71	1.02	1.10	0.99	1.10
PC_321 iso 2	1.14	0.94	0.75	0.84	0.93	0.88
PC_321 iso 3	1.16	1.11	0.80	0.92	1.00	1.00
PC_322 iso 3	0.91	0.76	0.76	0.84	1.52	1.65
PC_330_160_170	1.03	1.28	0.94	1.05	1.24	1.15
PC_331 iso 1	1.11	0.99	1.02	1.12	1.06	1.20
PC_331 iso 2	1.10	0.96	0.79	0.91	1.07	1.03
PC_341_160_181	0.92	0.94	1.11	1.10	1.07	1.03
PC_343 iso 1	0.80	0.82	0.82	0.86	1.34	1.31
PC_343 iso 2	0.73	0.52	0.76	0.80	1.59	1.58
PC_352 iso 1	1.01	0.93	1.01	1.06	1.04	1.11
PC_352 iso 2	1.06	0.92	0.85	0.76	1.03	1.05
PC_361_180_181	1.28	1.36	1.09	1.32	0.78	0.79
PC_362	1.01	0.89	1.00	1.10	0.91	0.87
PC_363 iso 1	0.91	0.69	0.81	0.87	1.18	1.12
PC_363 iso 2	1.02	0.77	0.70	0.78	0.91	0.94
PC_364	1.16	1.09	0.78	0.76	0.95	0.86
PC_381	1.16	1.31	1.04	1.20	0.99	0.99
PC_382	1.13	1.07	0.91	1.08	0.68	0.71
PC_383 iso 1	0.98	0.84	0.88	1.00	0.87	0.87
PC_383 iso 2	1.05	0.84	0.94	0.79	0.91	0.97
PC_383 iso 3	1.09	1.16	0.75	0.87	1.05	1.15
PC_384 iso 1	0.81	0.59	0.64	0.74	1.12	1.03
PC_384 iso 2	1.15	1.14	0.69	0.79	0.82	0.87
PC_385	0.98	0.87	0.69	0.73	0.86	0.80
PC_386	1.01	0.91	0.82	0.88	0.81	0.90
PC_403	1.16	1.28	0.69	0.78	0.93	0.98
PC_404 iso 1	0.84	0.70	0.90	0.70	0.96	1.01
PC_404 iso 2	1.09	1.07	0.66	0.78	0.80	0.87
PC_405 iso 1	0.85	0.76	0.63	0.70	0.99	1.03
PC_405 iso 2	1.09	1.02	0.66	0.71	0.86	0.83
PC_406_181_225	0.94	0.84	0.62	0.72	0.85	0.92
PC_407_181_226	1.01	0.89	0.58	0.62	0.86	0.81
PC_424	0.86	0.76	0.60	0.75	0.81	0.85

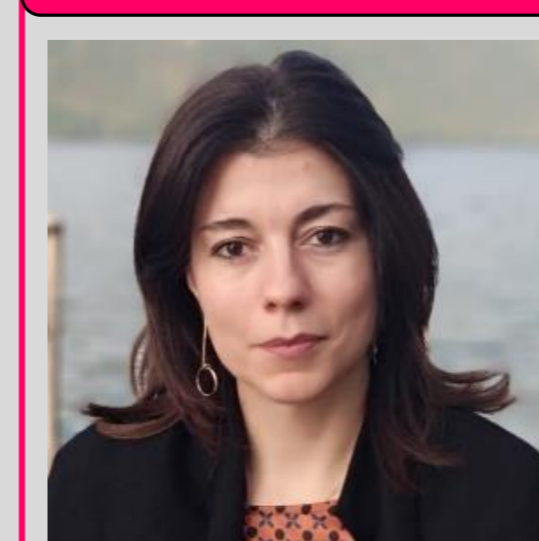
Lipidomic comparison of SH-SY5Y cells infected with HSV-1. (A) SH-SY5Y cells were infected with HSV-1 or left uninfected (MOCK) or treated with the indicated compounds or vehicle (DMSO), and then infected with HSV-1 (MOI of 1) and maintained in serum-free medium. At 24 and 48 hpi, total lipids from cells were extracted and analyzed by LC-MS/MS. The fold changes of lipid species peak area are visualized as a heat map showing the levels of upregulated (red, fold change > 1) and downregulated (blue, fold change < 1) glycerophospholipids. PC, glycerophosphocholine. n = 2 independent determinations. (B) Histogram reporting the total MS signal for PC in the different conditions.



CONCLUSIONS

- HSV-1 infection increases de novo fatty acid synthesis through transactivation of lipogenic enzymes.
- HSV-1 infection increased lipid storage.
- The anti-AD compounds CMS121 and C75 act inhibit HSV-1 replication.
- Some genes involved in AD are deregulated following HSV-1 infection.

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